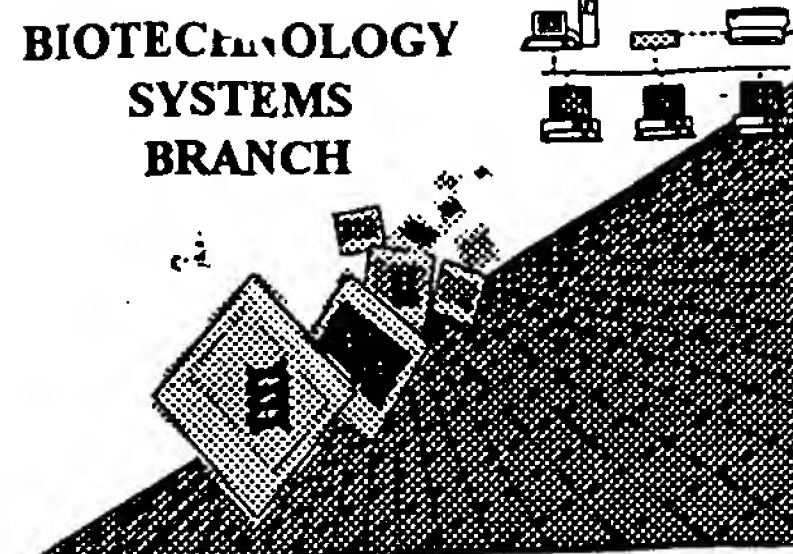


Shawin
Foley

RAW SEQUENCE LISTING **ERROR REPORT**



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/509,712
Source: 1648 RWA
Date Processed by STIC: 7/23/2001

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,**
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY**

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO).

Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:

<http://www.uspto.gov/web/offices/pac/checker>

Raw Sequence Listing Error Summary

ERROR DETECTED

SUGGESTED CORRECTION

SERIAL NUMBER: 09/509,712

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1 Wrapped Nucleics
 Wrapped Aminos The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
- 2 Invalid Line Length The rules require that a line not exceed 72 characters in length. This includes white spaces.
- 3 Misaligned Amino
 Numbering The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.
- 4 Non-ASCII The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
- 5 Variable Length Sequence(s) contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
- 6 PatentIn 2.0
 "bug" A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequence(s) . Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
- 7 Skipped Sequences
 (OLD RULES) Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence:
 (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
 (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 This sequence is intentionally skipped

 Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.
- 8 Skipped Sequences
 (NEW RULES) Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence.
 <210> sequence id number
 <400> sequence id number
 000
- 9 Use of n's or Xaa's
 (NEW RULES) Use of n's and/or Xaa's have been detected in the Sequence Listing.
 Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
 In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
- 10 Invalid <213>
 Response Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence
- 11 Use of <220> Sequence(s) missing the <220> "Feature" and associated numeric identifiers and responses.
 Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.
 (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
- 12 PatentIn 2.0
 "bug" Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.

1648

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/509,712

DATE: 07/23/2001

TIME: 15:47:30

Input Set : A:\W103960.txt

Output Set: N:\CRF3\07232001\I509712.raw

Does Not Comply
Corrected Diskette Needed

2 <110> APPLICANT: Rubin, Donald H.
3 Organ, Edward L.
4 DuBois, Raymond N.
6 <120> TITLE OF INVENTION: Mammalian Genes Involved in Viral
7 Infection and Tumor Suppression
9 <130> FILE REFERENCE: 01123.0004/P
11 <140> CURRENT APPLICATION NUMBER: US/09/509,712
11 <141> CURRENT FILING DATE: 2000-02-11
11 <150> PRIOR APPLICATION NUMBER: 60/062,021
12 <151> PRIOR FILING DATE: 1997-10-10
14 <160> NUMBER OF SEQ ID NOS: 127
16 <170> SOFTWARE: FastSEQ for Windows Version 3.0

This error type
exists in all sequences
(global error)

ERRORED SEQUENCES

18 <210> SEQ ID NO: 1
19 <211> LENGTH: 925
20 <212> TYPE: DNA
21 <213> ORGANISM: Rattus norvegicus
W--> 22 <400> SEQUENCE: 1
E--> 23 ggggggaaaac cnggnaattg ttttttgacg anccaaaaag ggggcnagna gcnnttttcc 60
E--> 24 tanatggggn cgggatcntn tccnaggana gattnatgga gtatnccttt ttgcnchnaa 120
E--> 25 ggttgattgc tcttgaaagg ntttgagggtg naattcctcc gtnagtttga ccgtagtcgg 180
E--> 26 atntgaagag ggattgttna gcagncataa tttcattccc tgnacacca gtaacnnttt 240
E--> 27 accgtcattt ggttggggaat tgatntcggg aggtancaan ggccacagtt atttattggt 300
E--> 28 ncggaggatt gcaccaattn ggccggctgc ctctganatc tgtttctcat ccatgccggt 360
E--> 29 tcaccagagac gaaagccgaa agcntcggga gtcctaactn tagtccttga aagtcattcc 420
30 cagctgcgta attgggctgt gcagagtcce agctcggtaa atatttgccc cgtgactgag 480
31 ctggagagaa tgctcctttc ttggtcctgg gcagctcttg gcagctcaca tgcactgttt 540
32 acctatcctc ccacattccc ccttgaggaa tcatcgtgcc tcggttccct taagtccctc 600
33 caacagaaaa caaggcagag tggaacgaag gaaagtgcgt ggccggttaga aagcctgtct 660
34 cgaatctgtc ccacgtgcct caggtagcgt tccaaacagc aaagattcta gtgaagaaaa 720
35 ataccgtccg gtcaattagt caggtggaca gagcaggacc cgggtgtctg gaagcctcgt 780
36 ccattcctct ggggaagggtg gggggggggcg tgtaatgcag ctctcaagaa gaaggatatt 840
37 ttgttttctc ggagaaactg ccattcccagg agctgagagt ggatcagtag gaaggcctgt 900
E--> 38 gacaggaagc agggaggttc agcng 925

see item 9 on Error Summary Sheet

see item 9

item 9

40 <210> SEQ ID NO: 2
41 <211> LENGTH: 554
42 <212> TYPE: DNA
43 <213> ORGANISM: Rattus norvegicus
W--> 44 <400> SEQUENCE: 2
E--> 45 caagatngan ggggcgggcg ttcgnccaga gagcgggtag ggaaggggaa ggcgccggtg 60
E--> 46 agccnggggtg cgganagcca gaccccaggc gtgggaaggg gagagagata gagcggccgg 120
E--> 47 ttgggaagag gaggaccgtg gttnataaat aacagaaagc ccagagggac gtanccatcc 180
E--> 48 gggatggaga gaggtaggga atccagntgt aagtcccaaa ctgccaccac cttcatnaga 240
E--> 49 actgcttcgt gtaagggtcac gcaccggggc agctgtccng agtggcggtc ctggcggtgt 300

item 9

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/509,712

DATE: 07/23/2001

TIME: 15:47:30

Input Set : A:\W103960.txt

Output Set: N:\CRF3\07232001\I509712.raw

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E--> 50 aagttagcta aagtnactgc aactccgnct gtgcagactg ntcgtaaatt ctctctgtcc 360
E--> 51 gccaaattct ccctcctatt aaacttttca cttcctttca cttagtttcc tnactttctt 420
E--> 52 caaacggaag ctgtaactga gcctgccacc cnganaentt gtgggttgcca tttttatgct 480
E--> 53 aaagtaatcg tgttttttat gcctgtcaac tcccttttca tntaaagcag ggcntaccct 540
54 attataactc tgcc 554
55 <210> SEQ ID NO: 3
56 <211> LENGTH: 891
57 <212> TYPE: DNA
58 <213> ORGANISM: Rattus norvegicus
W--> 59 <400> SEQUENCE: 3
E--> 60 ttngaaanaa tttccgtnaa ggtcngnaat nggccccgga aaaaatgngt tcctccccac 60
E--> 61 cttcattggn gcgatcctg ccngggaggg caatggttta acaaataatc tttnggagnt 120
E--> 62 ntggtngggg ggggagggac ncccacagan tcatgnggtg gttngggngg ngggcatcgt 180
E--> 63 tnngatatta tcacattntg ngaanctatg tnggggcttc ctttongaca ggtggtggtt 240
E--> 64 nnacangngg atgtgtgctt cttttttcag cagtgggtga cccggattct aagaccctta 300
E--> 65 cngtaacaat gccctntttt cetaagccta accagtcctt tangaggant gctcttggn 360
E--> 66 acccatgctg nntcacctag cettggntca catnttnnac acaggaaaag gcagcatgtc 420
E--> 67 ttntnggagc tcagcttatt cccttccent cccatccagn atctccctgg gntggatgag 480
E--> 68 gtggatgacg catcttcaaa gcacccacg tntcatggga tgtgcacagg agcttcgttg 540
E--> 69 gaaatgtgtt gcgcgaccag gcttgtgtag gaaacaacag actactcgaa attaaagtcn 600
E--> 70 taccttgacg ggttctcaga ggcttttaag cattaataaa catttgaatc ntaagaaggg 660
E--> 71 agcacagcat gtaatatnt tcaaattatc aggenttgca accttcatta gtttctctta 720
E--> 72 cgcagctggg ngtggtggtg tgtacctta atctcagcac tgaggaggca cngatatctc 780
E--> 73 catctctgtg acttcacagc cggcntcgcc agagcaagtt ccaggccacc cagatgagat 840
E--> 74 gctcacagag gggacctttt tntgatgacc aacgnagnat gcaagtaagg a 891
75 <210> SEQ ID NO: 4
76 <211> LENGTH: 974
77 <212> TYPE: DNA
78 <213> ORGANISM: Rattus norvegicus
W--> 79 <400> SEQUENCE: 4
E--> 80 aaaanaanat attccgnntc tnntagcnaa gaagtntnc gagenntccc ccgntttttt 60
E--> 81 aaaaaccnc ggaattccgn nntcgggntt taannngntt tttaanggcc cnaagnccn 120
E--> 82 nttattgccg ncntttcccc cccgctnttg cncctttta cttngagant ngtgntcna 180
E--> 83 agattttnaag gttnttgccc ccccggtttt tnttccctn nttttcccn nagnttttaa 240
E--> 84 accggtntgg gttncnantt nnttgnancc ncnattggg gtttccgntt accnggggtt 300
E--> 85 tcccccatgn ccgttccctc caatnttgna cttccnggt cngggtcna atnccnngna 360
E--> 86 acngntcnan ccttattgac aattaatttt tccttgngna ntctgncccc cngnantttg 420
E--> 87 gggttcttgg gngcagggcc tttttttent tggngcaan cncataaant ttaccagntt 480
E--> 88 gattgctaag gaagtancca tgggtgngaa ccccccttn ttntctccca gatggaacct 540
E--> 89 aggattttgg aactgcagag gcttcagggt cttgggaagc ggaggcaggn aaagattgga 600
E--> 90 gtgcactgtc cttttgcaat atgggggttg cctgcctgct ggctcntctc ctgctntntc 660
E--> 91 agatgggtgac tgaggctact tngcaggac tnggaataat catgtccagg tggctgccct 720
E--> 92 tccgagcaga aaggacaga cgtggggcga tgaagttgct atcgtttntt tttttttctg 780
93 cacagactgc aaagtgtgca gagggaggga ggctgtgcaa aaaaaaaaaa aaaaaaaaaa 840
94 aaaaaaaaaa ccgaggacgc agaagttaga ctgctgacct atttggtgca tgtgtgcca 900
E--> 95 tggaggaggg ggacctntt taaagggttc acgcggcacg cantgggnaa nngnncctnt 960
E--> 96 acgnnctcc caga 974
97 <210> SEQ ID NO: 5
98 <211> LENGTH: 850

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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/509,712

DATE: 07/23/2001

TIME: 15:47:30

Input Set : A:\W103960.txt

Output Set: N:\CRF3\07232001\I509712.raw

99 <212> TYPE: DNA

100 <213> ORGANISM: Rattus norvegicus

W--> 101 <400> SEQUENCE: 5

E--> 102	anttttccct	caagnaaant	ntggtttggg	caacttgaag	acgcttnnac	cnaaaaccct	60
E--> 103	tgnggagntt	ggngaccttn	ttaccgnaan	gagtgggaaa	cgttttcctc	cgggttnang	120
E--> 104	gttaggggga	cccgnnggaa	aatttttaaaa	ccnngngggc	tttttcgaat	taaggggaaa	180
E--> 105	ngcggtttng	gtnnntgaag	ggcggnggtt	tggagtcnna	gtccagagtt	gatttccacc	240
E--> 106	cacaaatntg	ggaggtgncg	gggaatgntg	ncnttttctt	gngatgaggg	ntgccgtnc	300
E--> 107	ggantaacag	ngnttgcntt	gtntngcnaa	acgaagagtn	tcctgnttgg	aataggngtt	360
E--> 108	cngttcgang	ganccagatt	tangngntgg	agnaaggatt	nggcagataa	angcntgaga	420
E--> 109	natgnancnt	ggancaggtc	nggnncnnagn	ntacagatga	tgnncccana	canganataa	480
E--> 110	ntncagatca	cagtcgtacc	cgnggctggg	ccatgaanag	ggcatcccca	gacnnacaca	540
E--> 111	ngccttnana	antgntcaga	gaaccancag	tggntanggg	ntgccnnnnn	naccagggaa	600
E--> 112	gacccggggc	gtgncggata	ttgacacanc	agatnncatt	tggggncggt	tcgagggttn	660
E--> 113	atgntcncg	agtacnagan	angatcntcc	aacccggaat	ncggtgctcc	ngtcgtccga	720
E--> 114	tgnaatgagt	cgncggnaa	cctcatatcc	aagaaacnat	acagcagtgg	nntccgagtc	780
E--> 115	tcgtatantc	nttgcgggng	gaggctatnt	tcagaggnc	agattaccgt	tagcgggana	840
E--> 116	aagtngaana						850

117 <210> SEQ ID NO: 6

118 <211> LENGTH: 531

119 <212> TYPE: DNA

120 <213> ORGANISM: Rattus norvegicus

W--> 121 <400> SEQUENCE: 6

E--> 122	ttgnggcngg	gtctcctctg	ngtgngngtn	tcccnanag	gggggggtctc	acagtgtng	60
E--> 123	ngtctnntgt	ctgtgtngtg	cccctgtccn	catctctcac	nccagggaga	gagatgtgag	120
E--> 124	ananacatca	gagatctctn	gnacagtgtt	tcacaagagt	ctatcnana	gagcacatct	180
E--> 125	gcccggggng	anacacaact	ctaaatgtgt	ctcanntgat	ctctctnttg	tgtctctnac	240
E--> 126	atatngggac	atgtctctcag	agtatnggnt	ctcttgngcn	ctnttgacac	cacacacaca	300
E--> 127	cacacacaca	cacacacaca	cacncttctc	tctggcacag	ggntatggca	nagcacatnt	360
E--> 128	tnngagntca	nagctntata	tgagtgtgtg	gcgaaaggng	tnatnanann	gacnncccca	420
E--> 129	gcnnatatag	gggggngnnc	tctngggctc	tcttnggnaa	tntgngggng	agtctgcnc	480
E--> 130	cacaggcgct	cnnacccanc	nnnttggggc	ccccaggng	tttttcnccc	c	531

131 <210> SEQ ID NO: 7

132 <211> LENGTH: 572

133 <212> TYPE: DNA

134 <213> ORGANISM: Rattus norvegicus

W--> 135 <400> SEQUENCE: 7

E--> 136	ttttnttgtg	gccctttaaa	ctctgngtgn	ccgtntnccc	nagagggggg	gtctcacaag	60
E--> 137	gagacancgg	nnacacagag	gttttgngnn	tattgngagt	ctctgcgcac	nccananttt	120
E--> 138	aaccncgggg	ntcntgtttt	tatttttaaaa	aaaaagagtc	ncatgtntat	ttctctnatg	180
E--> 139	tgaaaatcnc	attcanagtt	ntgggggtttc	ccntgaggag	anatagagtt	tcacactctt	240
E--> 140	ctctccgagg	ggtentcnc	tgtntctccc	caatgtgngn	ggnacacaca	tgngggcccn	300
E--> 141	agggggtgng	ctctctctgc	ncagggcnc	ccccaanang	tagaganaca	ntgtgggtgt	360
E--> 142	tcacaacaca	attcncgaga	nattntgttc	cncantggnn	gtctnagntc	ncatgttgtg	420
E--> 143	gngacangtt	agnncncccc	atnttncccc	ccctttcaca	ctgccccnag	agagagaaan	480
E--> 144	tctnggcccc	ctctanannt	nttttttaaat	cnccccnnac	cacaggtnnt	cccagggtat	540
E--> 145	gngacntcnc	cnnccccncn	aaagatntgc	nc			572

146 <210> SEQ ID NO: 8

147 <211> LENGTH: 906

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/509,712

DATE: 07/23/2001

TIME: 15:47:30

Input Set : A:\W103960.txt

Output Set: N:\CRF3\07232001\I509712.raw

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148 <212> TYPE: DNA
149 <213> ORGANISM: Rattus norvegicus
W--> 150 <400> SEQUENCE: 8
E--> 151  tgggagtcctc tctcatatgg cgcnttcncc aaaggggngt ctctntccng agncgcanac      60
E--> 152  gcgagaanac tctgtnnant ngctctcccc cncncnaca gngtganant caaaacctct      120
E--> 153  agagccccc agaaancccc tntctcaaan aaagagaaag agaagancga gnagnagaga      180
E--> 154  ganaganaga gagagagtgt ggancntntt cctcngancc ccannnanan ngtgnggcnc      240
E--> 155  actcncnngt gnnngnacc ccnggggatt tncgcgtgtc cccttgngct ctgtntanga      300
E--> 156  gananatatg tntagtctct ctntcgcgcc ctccngtgtc acgtgtgcgg ggcccnngag      360
E--> 157  acacagacac ntctctcang gggaacacat anngactcnc acntgtgttt atattcnccc      420
E--> 158  ctcccnctca cacanacaca cacacagnag atattnngct actctctctc tgtcacaggg      480
E--> 159  gtacanattt antctnggcc anaccctctc cngaagngng ggcanngtaa accccgcccc      540
E--> 160  ctctcngaga angngagggc gntttacntt ccngtgggcg tgtncngncc cccgagactc      600
E--> 161  cccttngnac cccctntna accctctntt tgaacncaac ncacctccc cnttttctcg      660
E--> 162  gggngngncc ngcncccnct ctcncaaaaa aaattnnaan ttngtccctt ncccnctntt      720
E--> 163  ttcnggnana aaccgtgtcc ggggggggan nactcttttt tgnccctaaa atcaantttt      780
E--> 164  ttcccttttt ccnggggacc cccgnnttcc tttttaaaaa aaaanaaccc tttctccctt      840
E--> 165  ttaaaagnac ccnttttttc naaaaccgtt ccgnatttaa ttcctaaatt cccttcccn      900
E--> 166  ncccg      906
167 <210> SEQ ID NO: 9
168 <211> LENGTH: 914
169 <212> TYPE: DNA
170 <213> ORGANISM: Rattus norvegicus
W--> 171 <400> SEQUENCE: 9
E--> 172  gggatgngcc ctcagatcaa tacaccctc ngggggngtc tctctctatc tccncagna      60
E--> 173  gactcccatc tctntntntn ccccaganc tggngaacgg nggtgtgnga nccntntctg      120
E--> 174  ttctcnantc tctaaaagng cnaaaagcgc ananacacgn gcctctctat anatctcacg      180
E--> 175  tgtcccnngn nctctcngac ccctnntctg tntgagagac accctntctc aaaatatagt      240
E--> 176  gtacacgngc tttgnggctc tcccttttct tctccactnt tgagnngaa acgcgnggtt      300
E--> 177  ntctctgaga tgtaganagn gtccctnct cnatatatgt gttncctact ccnaggngng      360
E--> 178  tctcataaaa atcnctntct tcaacaccac cncctcnacc cccncacga gaacacntcn      420
E--> 179  ccaccnncan gacacaaana naaggngtnn anaacccan aaaaactnng ntntcngntt      480
E--> 180  tacacacaca cacacncacn ctcnncaca cccccacna aatgggagaa aaaacagaga      540
E--> 181  ggngtggttg ttngntcaa caccntntta cctctctgnt gnnanttgag aaaatatctc      600
E--> 182  tntncttacc cctctccctt ctctgtgtgt ngannatatc ngntctagat gtcctnacc      660
E--> 183  tcccaaacc tttctcnggn agagacntct ctntnttttt ccccncttc catttgaaan      720
E--> 184  anangagaag gnccaaaaag gngggngtct tctcggaat ncnccctttt ggccccccaa      780
E--> 185  cctgggtttt tttccctt ccttttaatn antttttcna nacaanctt tnngngtttn      840
E--> 186  ggaaaangcc tttnnctggn nnttttttcc cttcccttt tnnangggnt tcccccccc      900
E--> 187  ccngaatttt tttt      914
200 <210> SEQ ID NO: 11
201 <211> LENGTH: 880
202 <212> TYPE: DNA
203 <213> ORGANISM: Rattus norvegicus
W--> 204 <400> SEQUENCE: 11
E--> 205  acccaatctt nanggtggca gtgnggnga tcttaacggt ttttnagaaa aaaaantnct      60
E--> 206  tcgctcncac cccaagcct cccnttctta ncagcttttt tatangaaa aagatgataa      120
E--> 207  cgaaatttta aaaaccgtcg ttagaggaaa tgaaggttca gccgaccatt acctganagt      180
E--> 208  aatgaaggtn ttccggaggg ttgccttcca atcccagatg gatttgagtt tcaggatcaa      240

```

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/509,712

DATE: 07/23/2001

TIME: 15:47:30

Input Set : A:\W103960.txt

Output Set: N:\CRF3\07232001\I509712.raw

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E--> 209   ttcagttacc gntgaccatc caccnncctc cngtataatc attngatgag gatgaatggt      300
E--> 210   gagtgagtga tgatgatgat gatgatgatg aagggatgag aagnacacta tgataacaag      360
E--> 211   tgtctcagtc cacattaagg tttgcctgna aattagtgca taagccatgg gagacaaatt      420
E--> 212   cttttcnnac acaattaata gtntcttant ccttcccatc ttctctgccc cattctgttt      480
      213   tccaccacag gtctgcagcg ggctacagct tccagtctcc aagcaaatac cagaactgga      540
      214   ggagaaaatt ccagtcacagt gagtcatggg cagggggagg ggtggggtaa gggcagtggc      600
E--> 215   gctcattcct nacatgggtg cttctcttgc cttagcctggg atctgagggc aagagaacct      660
      216   gtaagcttga tttgatttcc actgctgact ggagtcactg ccaagggatt tgggacttct      720
      217   ccctctctct ctctaacctg aaatccttag gattctatta tttcaccgga ccagagctgt      780
E--> 218   agcagagatg agctccaagt ttgaaatgag aaaggggaaa ttgagagcta tgagctaggn      840
E--> 219   gcgaaagncc ccacaaagnn tttggcaagt agaaaagncg      880
      220   <210> SEQ ID NO: 12
      221   <211> LENGTH: 909
      222   <212> TYPE: DNA
      223   <213> ORGANISM: Rattus norvegicus
W--> 224   <400> SEQUENCE: 12
E--> 225   cgngagnngg cagggannna gnggggagcn ngagaggaga aggagaaggn nnggnaggng      60
E--> 226   nngngagnaa cgggcgggan cnnngacga gagaangggg aggggancga agngcggngg      120
E--> 227   nagacggtgc nnggggggga ggggcaggag nggnagagag gcangagnng agnggggaca      180
E--> 228   agcnaaaanc gaggaggnan gangngangg nngngngnc gaaggcgcn aagnnggtcg      240
E--> 229   gngagcggna gnggnnaaac tggggaacga gacagacggc ccnncggng gcangnggga      300
E--> 230   gagnnncgcc agngagagna gncagnanca gancanggga ggggggggan ncacnggcgg      360
E--> 231   gagggncgan gacggnnngn annngnnaga ggcannnnnc gccnanagn ngagngagg      420
E--> 232   cangagtgn cgnngagnag acaggcccgc gcncggggg cagacnnng ncaccaccga      480
E--> 233   ggggtggngg ggcncggaga naagaccaga ggnngagggg cganggcnn ggtnngcccg      540
E--> 234   ggccnccna aaaaaanncc gaaaaaaan aaggggcgcn gcngggcngg ggaggagcgc      600
E--> 235   ntncgtang tngantgacg gaggccngna atngggccgn gccanncnag ggcgnagagg      660
E--> 236   cccaagncg gnagngnaa gnanagancc ngnggtngg gagnganagn gcnnngncc      720
E--> 237   naccnccngn gttganggn cccacgncgg ngcaggccgn nnaaagnag tcccnaaaa      780
E--> 238   nntcnggtn tnacancgnc ccggggncgc cgcngngtcc cgncacacng gannncggag      840
E--> 239   anngcctnnt ntctncacan ggngccanac nngntgctat gcaaaagggg cgnacttcna      900
E--> 240   gaaaaagnc      909
      241   <210> SEQ ID NO: 13
      242   <211> LENGTH: 927
      243   <212> TYPE: DNA
      244   <213> ORGANISM: Rattus norvegicus
W--> 245   <400> SEQUENCE: 13
E--> 246   cctttattcg gaggcaggga nnncttgtcc gggaangtta aacgtttttt aaaagggggg      60
E--> 247   nccnnggggg gggggnntnt ccagggaant aaaanggtgn gttggggggg aaaaatttat      120
E--> 248   tttnaaaaag ggcgncnat ataaangacn ttcggggggg tttgaanagg gccggaancn      180
E--> 249   tcgacgggtt tccggngggg ganaaggana aggnacgca cgggatttct tnccttttt      240
E--> 250   tngcaaattg cngcaggana ccaccgggtg gggnggtttt gttttccgtn aagaaagcgg      300
E--> 251   gngtggaata acangataa acgggaagan ggggttatnt nggttagnaa ttgnttccag      360
E--> 252   ngngccagg aaattggcct gtccaaaatt cttttccng cttttaagac aggcagggtat      420
E--> 253   tatttgagc caggttatta cnataggnaa gtaaataaca atgggtaagt gcctggcaca      480
E--> 254   ggccagggtg agtagggcat gtatggaatg ttaaacatta cccttcatcc tgagaaanaa      540
E--> 255   aanacaagna anaaaggctg gtctcacata tcccaaagct ttatcttct aggtgcccc      600
E--> 256   tggtgaacgt taagccaagc ntatgantca caagggacga catgggcagg ntagggtaca      660
E--> 257   gaatcagtn tcagagactc caggggcacc cctgattccc tttgctgtca cacagacact      720

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FYI →

Use of n and/or Xaa has been detected in the Sequence Listing. Review the Sequence Listing to insure a corresponding explanation is presented in the <220> to <223> fields of each sequence using n or Xaa.